	r: 09/6	n-ASCII to ASCII			dited by:	( ) _
	10年表		· 'E		r en Skrafaggy Grand bereit	
Chang	ed the margins in	cases where the	sequence text wa	is was ped o	down to the ne	xt line.
Edited	a format error in	the Current Applica	ation Data sectio	n, specifically:	**********	·, .
Edited applicate	the Current Appli nt was  the p	ication Data section nor application data	n with the actual a; or  other _	current numbe	er. The numbe	er inputte
Added	he mandatory he	eading and subhea	dings for "Currer	t Application [	Data".	
Edited t	he "Number of S	Sequences" field. T	he applicant spe	lled out a num	ber instead of	using ar
Change	d the spelling of	a mandatory field (	the headings or	subheadings),	specifically:	
Correcte	ed the SEQ ID N	O when obviously i	ncorrect. The se	quence numb	ers that were	edited w
Inserted	or corrected a n	ucleic number at th	e end of a nuclei	c line. SEQ I	D NO's edited	•
Correcte	d subheading pla	acement. All response below the subh	enses must be on	the same line	as each subh	eading.
		nse below the subh	ieauny, uns was	moved to its a	ihbiobuate bla	CO.
Inserted						
	colons after hea	adings/subheadings	s. Headings edite	ed included:		
	· · · · · · · · · · · · · · · · · · ·	adings/subheadings eadings used by an				
Deleted	extra, invalid, he		applicant, specif	ically:	tary initials/file	name at
Deleted Deleted	extra, invalid, he non-ASCII	eadings used by an garbage" at the be ughout text;	applicant, specif	ically:	tary initials/file	name at
Deleted Deleted pag	extra, invalid, he long to mandatory head	eadings used by an	applicant, specif ginning/end of fil er invalid text, su	ically:	tary initials/file	name at
Deleted  Deleted  pag  Inserted  Correcte	extra, invalid, he non-ASCII non-ASCII numbers throu mandatory head d an obvious err	radings used by an garbage at the be ughout text;  oth	applicant, specifinging ginning/end of filer invalid text, su	es; Secre		name at
Deleted Deleted pag Inserted Correcte	extra, invalid, he formation in the form	garbage" at the be ughout text;  oth dings, specifically:	applicant, specification of filer invalid text, subspecifically:	es;  secre		name at
Deleted  Deleted  pag  Inserted  Correcte  Edited id	extra, invalid, he long to mandatory head an obvious error in the	garbage" at the be ughout text; oth dings, specifically: or in the response, upper case is used	applicant, specification of filer invalid text, subspecifically:  but lower case is access field, specification of the properties of the p	es;  secre	ice versa.	) ( )
Deleted  Deleted  pag Inserted Correcte Edited id Correcte A *Hard I	extra, invalid, he non-ASCII and an obvious ementifiers where under an error in the page Break code and ing stop code	garbage" at the be ughout text;  oth dings, specifically: or in the response, upper case is used Number of Sequence was inserted by the manino acid second and second second or in amino acid second o	applicant, specifically:  but lower case is need field, specifically:  the applicant. All quences and adiaset.	es;  secretich as required, or vocally:	ice versa.	led.
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Deleted  Deleted  pag Inserted Correcte  Edited id Correcte  A "Hard I	extra, invalid, he non-ASCII and an obvious ementifiers where under an error in the page Break code and ing stop code	garbage" at the be ughout text;  oth dings, specifically: or in the response, upper case is used. Number of Sequences corrected by the continuation acid sequences corrected.	applicant, specifically:  but lower case is need field, specifically:  the applicant. All quences and adiaset.	es;  secretich as required, or vocally:	ice versa.	led.
Deleted  Deleted  pag Inserted Correcte Edited id Correcte A *Hard I Deleted ex due to a P Other:	extra, invalid, he non-ASCII and an obvious ementifiers where under an error in the page Break code and ing stop code	garbage" at the be ughout text;  oth dings, specifically: or in the response, upper case is used. Number of Sequences corrected by the continuation acid sequences corrected.	applicant, specifically:  but lower case is aces field, specifically: the applicant. All quences and adjustically:	es;  secretich as required, or vocally:	ice versa.	led.

P4/09

RAW SEQUENCE LISTING

4 <110> APPLICANT: LAMBERTY, MIREILLE

DATE: 06/17/2002

PATENT APPLICATION: US/09/673,274

TIME: 19:40:18

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF3\06172002\I673274.raw

BULET, PHILLIPE BROOKHART, GARY 6 7 HOFFMAN, JULES 9 <120> TITLE OF INVENTION: GENE CODING FOR HELIOMICINE, AND USE THEREOF 12 <130> FILE REFERENCE: A33595-PCT-USA 14 <140> CURRENT APPLICATION NUMBER: 09/673,274 15 <141> CURRENT FILING DATE: 1999-04-12 17 <150> PRIOR APPLICATION NUMBER: PCT/FR99/00843 18 <151> PRIOR FILING DATE: 1999-04-12 20 <150> PRIOR APPLICATION NUMBER: FR 98 04933 21 <151> PRIOR FILING DATE: 1998-04-15 23 <160> NUMBER OF SEQ ID NOS: 38 25 <170> SOFTWARE: FastSEQ for Windows Version 4.0 27 <210> SEQ ID NO: 1 28 <211> LENGTH: 147 29 <212> TYPE: DNA 30 <213> ORGANISM: Artificial Sequence 32 <220> FEATURE: 33 <223> OTHER INFORMATION: SYNTHETIC POLYNUCLEOTIDE 36 <400> SEQUENCE: 1 37 agcttggata aaagagacaa gttgattggc agctgtgttt ggggcgccgt caactacact 60 38 agtgactgca acggcgagtg caagcgccgc ggttacaagg gtggccattg tggatccttc 120 39 gctaacgtta actgttggtg tgaaacc 41 <210> SEQ ID NO: 2 42 <211> LENGTH: 169 43 <212> TYPE: DNA 44 <213> ORGANISM: Artificial Sequence 46 <220> FEATURE: 47 <223> OTHER INFORMATION: SYNTHETIC POLYNUCLEOTIDE 50 <400> SEQUENCE: 2 51 gataagetta teggtteetg egtgtggggt getgtgaaet acaetteega ttgcaaeggt 60 52 gagtgcaaga ggagggtta caagggtggt cactgcggtt ccttcgctaa cgtgaactgc 120 53 tggtgcgaga cttgagagct cggcgaggcg aacgtgtcga cggatccgg 55 <210> SEQ ID NO: 3 56 <211> LENGTH: 261 57 <212> TYPE: DNA 58 <213> ORGANISM: Artificial Sequence 60 <220> FEATURE: 61 <223> OTHER INFORMATION: SYNTHETIC POLYNUCLEOTIDE 64 <400> SEQUENCE: 3 65 ccatgggttt cgtgcttttc tctcagcttc catctttcct tcttgtgtct actcttcttc 60

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF3\06172002\I673274.raw

66 ttttccttgt gatctctcac tcttgccgtg ccgataagct tatcggttcc tgcgtgtggg 120 67 gtgctgtgaa ctacacttcc gattgcaacg gtgagtgcaa gaggaggggt tacaagggtg 180 68 gtcactgcgg ttccttcgct aacgtgaact gctggtgcga gacttgagag ctcggcgagg 240 69 cgaacgtgtc gacggatccg g 71 <210> SEQ ID NO: 4 72 <211> LENGTH: 120 73 <212> TYPE: DNA 74 <213> ORGANISM: Artificial Sequence 76 <220> FEATURE: 77 <223> OTHER INFORMATION: SYNTHETIC POLYNUCLEOTIDE 80 <400> SEQUENCE: 4 81 gcgtcgacgc gatgggtttc gtgcttttct ctcagcttcc atctttcctt cttgtgtcta 60 82 ctcttcttct tttccttgtg atctctcact cttgccgtgc tggagacgcg aattcacaca 120 85 <210> SEQ ID NO: 5 86 <211> LENGTH: 75 87 <212> TYPE: DNA 88 <213> ORGANISM: Artificial Sequence 90 <220> FEATURE: 91 <223> OTHER INFORMATION: SYNTHETIC POLYNUCLEOTIDE 94 <400> SEQUENCE: 5 95 gcgtcgacgc gatgggtttc gtgcttttct ctcagcttcc atctttcctt cttgtgtcta 60 96 ctcttcttct tttcc 98 <210> SEQ ID NO: 6 99 <211> LENGTH: 72 100 <212> TYPE: DNA 101 <213> ORGANISM: Artificial Sequence 103 <220> FEATURE: 104 <223> OTHER INFORMATION: SYNTHETIC POLYNUCLEOTIDE 107 <400> SEQUENCE: 6 108 tcgccggcac ggcaagagta agagatcaca aggaaaagaa gaagagtaga cacaagaagg 60 109 aaagatggaa gc 111 <210> SEQ ID NO: 7 112 <211> LENGTH: 80 113 <212> TYPE: DNA 114 <213> ORGANISM: Artificial Sequence 116 <220> FEATURE: 117 <223> OTHER INFORMATION: SYNTHETIC POLYNUCLEOTIDE 120 <400> SEQUENCE: 7 121 gataagetta teggtteetg egtgtggggt getgtgaaet acaetteega ttgcaaeggt 60 122 gagtgcaaga ggaggggtta 124 <210> SEQ ID NO: 8 125 <211> LENGTH: 109 126 <212> TYPE: DNA 127 <213> ORGANISM: Artificial Sequence 129 <220> FEATURE: 130 <223> OTHER INFORMATION: SYNTHETIC POLYNUCLEOTIDE 133 <400> SEQUENCE: 8 134 ccggatccgt cgacacgttc gcctcgccga gctctcaagt ctcgcaccag cagttcacgt 60 135 tagcgaagga accgcagtga ccaccettgt aacccctcct ettgcacte

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF3\06172002\1673274.raw

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204 <213> ORGANISM: Artificial Sequence

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF3\06172002\I673274.raw

207 <223> OTHER INFORMATION: SYNTHETIC POLYNUCLEOTIDE 210 <400> SEQUENCE: 14 211 ccgaattcgg cgccgtgcac gatgcagaag agcacgagga ggaagagggc 50 213 <210> SEQ ID NO: 15 214 <211> LENGTH: 81 215 <212> TYPE: DNA 216 <213> ORGANISM: Artificial Sequence 218 <220> FEATURE: 219 <223> OTHER INFORMATION: SYNTHETIC POLYNUCLEOTIDE 222 <400> SEQUENCE: 15 223 totagaatgg cotgoaccaa caacgocatg agggocotot tootootoot gotottotgc 60 224 atcgtgcacg gcgccgaatt c 226 <210> SEQ ID NO: 16 227 <211> LENGTH: 24 228 <212> TYPE: DNA 229 <213> ORGANISM: Artificial Sequence 231 <220> FEATURE: 232 <223> OTHER INFORMATION: SYNTHETIC POLYNUCLEOTIDE 235 <400> SEQUENCE: 16 24 236 gataagetta teggtteetg egtg 238 <210> SEQ ID NO: 17 239 <211> LENGTH: 32 240 <212> TYPE: DNA 241 <213> ORGANISM: Artificial Sequence 243 <220> FEATURE: 244 <223> OTHER INFORMATION: SYNTHETIC POLYNUCLEOTIDE 247 <400> SEQUENCE: 17 32 248 ggctcgagtc aagtctcgca ccagcagttc ac 250 <210> SEQ ID NO: 18 251 <211> LENGTH: 213 252 <212> TYPE: DNA 253 <213> ORGANISM: Artificial Sequence 255 <220> FEATURE: 256 <223> OTHER INFORMATION: SYNTHETIC POLYNUCLEOTIDE 259 <400> SEQUENCE: 18 260 totagaatgg cotgoaccaa caacgocatg agggoodtot tootootoot gotottotgo 60 261 atcgtgcacg gcgataagct tatcggttcc tgcgtgtggg gtgctgtgaa ctacacttcc 120 262 gattgcaacg gtgagtgcaa gaggagggt tacaagggtg gtcactgcgg ttccttcgct 180 263 aacgtgaact gctggtgcga gacttgactc gag 265 <210> SEQ ID NO: 19 266 <211> LENGTH: 838 267 <212> TYPE: DNA 268 <213> ORGANISM: Artificial Sequence 270 <220> FEATURE: 271 <223> OTHER INFORMATION: SYNTHETIC POLYNUCLEOTIDE W--> 274 <221> NAME/KEY: promoter 275 <222> LOCATION: (7)...(532)

W--> 277 <221> misc structure

278 <222> LOCATION: (533)...(568)

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF3\06172002\1673274.raw

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     281 <222> LOCATION: (569)...(832)
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     285 actatggaag tattatgtga geteageaag aageagatea atatgeggea eatatgeaae 120
     286 ctatgttcaa aaatgaagaa tgtacagata caagatccta tactgccaga atacgaagaa 180
     287 qaatacqtaq aaattqaaaa aqaaqaacca qqcqaaqaaa aqaatcttqa aqacqtaaqc 240
     288 actgacgaca acaatgaaaa gaagaagata aggtcggtga ttgtgaaaga gacatagagg 300
     289 acacatgtaa ggtggaaaat gtaagggcgg aaagtaacct tatcacaaag gaatcttatc 360
     290 ccccactact tatcctttta tatttttccg tgtcattttt gcccttgagt tttcctatat 420
     291 aaggaaccaa gttcggcatt tgtgaaaaca agaaaaaatt tggtgtaagc tattttcttt 480
     292 gaagtactga ggatacaact tcagagaaat ttgtaagttt gtagatctcg attctagaag 540
     293 gootgaatto gagotoggta coggatocaa ttoocgatog ttoaaacatt tggcaataaa 600
     294 gtttcttaag attgaatcct gttgccggtc ttgcgatgat tatcatataa tttctgttga 660
    295 attacgttaa gcatgtaata attaacatgt aatgcatgac gttatttatg agatgggttt 720
    296 ttatgattag agtcccgcaa ttatacattt aatacgcgat agaaaacaaa atatagcgcg 780
    297 caaactagga taaattatcg cgcgcggtgt catctatgtt actagatcgg ggatcgat
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     300 <211> LENGTH: 1036
     301 <212> TYPE: DNA
     302 <213> ORGANISM: Artificial Sequence
     304 <220> FEATURE:
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W--> 308 <221> NAME/KEY: promoter
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     320 ctatgttcaa aaatgaagaa tgtacagata caagatccta tactgccaga atacgaagaa 180
     321 gaatacqtaq aaattqaaaa agaaqaacca qqcqaaqaaa agaatcttga agacgtaagc 240
     322 actgacgaca acaatgaaaa gaagaagata aggtcggtga ttgtgaaaga gacatagagg 300
     323 acacatgtaa ggtggaaaat gtaagggcgg aaagtaacct tatcacaaag gaatcttatc 360
     324 ccccactact tatcctttta tatttttccg tgtcattttt gcccttgagt tttcctatat 420
     325 aaggaaccaa gttcggcatt tgtgaaaaca agaaaaaatt tggtgtaagc tattttcttt 480
     326 gaagtactga ggatacaact tcagagaaat ttgtaagttt gtagatctcg attctaga
                                                                            586
     327 atg gcc tgc acc aac acc gcc atg agg gcc ctc ttc ctc ctc gtg ctc
     328 Met Ala Cys Thr Asn Asn Ala Met Arg Ala Leu Phe Leu Leu Val Leu
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     331 ttc tgc atc gtg cac ggc gat aag ctt atc ggt tcc tgc gtg tgg ggt
                                                                            634
     332 Phe Cys Ile Val His Gly Asp Lys Leu Ile Gly Ser Cys Val Trp Gly
                      20
     335 gct gtg aac tac act tcc gat tgc aac ggt gag tgc aag agg agg ggt
                                                                            682
     336 Ala Val Asn Tyr Thr Ser Asp Cys Asn Gly Glu Cys Lys Arg Arg Gly
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                  35
                                      40
                                                                            730
     339 tac aag ggt ggt cac tgc ggt tcc ttc gct aac gtg aac tgc tgg tgc
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Does Not Comply

Corrected Diskette Needer

PCT09

RAW SEQUENCE LISTING

DATE: 06/05/2002

PATENT APPLICATION: US/09/673,274

TIME: 17:08:44

Input Set : A:\A33595-PCT-USA sequence listing.txt

Output Set: N:\CRF3\06052002\1673274.raw

4 <110> APPLICANT: LAMBERTY, MIREILLE

BULET, PHILLIPE

BROOKHART, GARY 6

HOFFMAN, JULES 7

9 <120> TITLE OF INVENTION: GENE CODING FOR HELIOMICINE, AND USE

THEREOF 10

12 <130> FILE REFERENCE: A33595-PCT-USA

14 <140> CURRENT APPLICATION NUMBER: 09/673,274

C--> 15 <141> CURRENT FILING DATE: 2001-12-18

17 <150> PRIOR APPLICATION NUMBER: PCT/FR99/00843

18 <151> PRIOR FILING DATE: 1999-04-12

20 <150> PRIOR APPLICATION NUMBER: FR 98 04933

21 <151> PRIOR FILING DATE: 1998-04-15

23 <160> NUMBER OF SEQ ID NOS: 38

25 <170> SOFTWARE: FastSEQ for Windows Version 4.0

## ERRORED SEQUENCES

557 <210> SEQ ID NO: 38

558 <211> LENGTH: 26

559 <212> TYPE: DNA

560 <213> ORGANISM: Artificial Sequence

562 <220> FEATURE:

563 <223> OTHER INFORMATION: SYNTHETIC POLYNUCLEOTIDE

566 <400> SEQUENCE: 38

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26

VARIABLE LOCATION SUMMARY

PATENT APPLICATION: US/09/673,274

DATE: 06/05/2002 TIME: 17:08:45

Input Set : A:\A33595-PCT-USA sequence listing.txt

Output Set: N:\CRF3\06052002\1673274.raw

## Use of n's or Xaa's (NEW RULES):

Use of n's and/or Xaa's have been detected in the Sequence Listing.
Use of <220> to <223> is MANDATORY if n's or Xaa's are present.
in <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

Seq#:38; N Pos. 27